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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/684,554

DATE: 10/19/2000
TIME: 15:08:53

Input Set : A:\875-024US1.txt.txt
Output Set: N:\CRF3\10192000\I684554.raw

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4 <110> APPLICANT: Engelhardt, John F.
5      Duan, Dongsheng
6      Yan, Ziyi
8 <120> TITLE OF INVENTION: Adeno-associated virus vectors and uses thereof
10 <130> FILE REFERENCE: 875.024US1
C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/684,554
C--> 12 <141> CURRENT FILING DATE: 2000-10-06
12 <150> PRIOR APPLICATION NUMBER: US 60/158,209
13 <151> PRIOR FILING DATE: 1999-10-07
15 <160> NUMBER OF SEQ ID NOS: 13
17 <170> SOFTWARE: FastSEQ for Windows Version 4.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 20
21 <212> TYPE: DNA
22 <213> ORGANISM: Adeno-associated virus
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27 <210> SEQ ID NO: 2
28 <211> LENGTH: 19
29 <212> TYPE: DNA
30 <213> ORGANISM: Adeno-associated virus
32 <400> SEQUENCE: 2
33 gggcgagcc tatggaaaa
35 <210> SEQ ID NO: 3
36 <211> LENGTH: 505
37 <212> TYPE: DNA
38 <213> ORGANISM: Artificial Sequence
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40 <223> OTHER INFORMATION: A consensus sequence
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43 cgggggtcgt tgggggtca gccaggcgccatattaccg taagttatgt aacgactgca 60
44 ggcataaag ctgcgaatca tcggtagata agtagcatgg cgggttaatc attaactaca 120
45 aggaaccctt agtgatggag ttggccactc cctctctgcg cgctcgtctcg ctgcgtgagg 180
46 ccgggcgacc aaaggtcgcc cgacggcccg gctttgcgg ggcggcctca gtgagcgagc 240
47 gagcgcgcag ctgcgcgttc gtcgcgtcac tgaggccgc cgggaaagc ccggcggtcg 300
48 ggcacettt ggtcgccccgg cctcagcgag cgagcgagcg cgcagagagg gagtgccaa 360
49 ctccatcaact aggggttct ttagttaat gattaacccg ccatgctact tatctacagc 420
50 ttgcatgcat gtgagccaaa ggcagccaaa aggcaggaa ccgtaaaaag gccgcgttc 480
51 tggcggtttt ccataaggctt cgcggcc 505
53 <210> SEQ ID NO: 4
54 <211> LENGTH: 272
55 <212> TYPE: DNA
56 <213> ORGANISM: AAV circular intermediate, clone p81
58 <400> SEQUENCE: 4
59 gcatgcaagc ttagataag tagcatggcg ggtaatcat taactacaag gaacccttag 60
60 ttagggattt ggccactccc tctctgcgcg ctgcgtcgct cactgaggcc gggcgccaa 120
61 aggtcgcccg acgccccggc ttgtccccggg cggcctcagt gagcgagcga gcgcgcagag 180

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62	agggagtggc caactccatc actaggggtt ccttgttagtt aatgattaac ccgcctatgc	240
63	acttatctac cgatgaattc gagcttgcac gc	272
65	<210> SEQ ID NO: 5	
66	<211> LENGTH: 300	
67	<212> TYPE: DNA	
68	<213> ORGANISM: AAV circular intermediate, clone p79	
70	<400> SEQUENCE: 5	
71	gcatcaagc ttagataag tagcatggcg ggttaatcat taactacaag gaacccttag	60
72	ttagggatt ggccactccc tctctgcgcg ctgcgtcgct cactgaggcc gggcgccgc	120
73	tcgctcgctc actgaggccg ggcgacaaa ggtcgccgaa gccgggctt tgccccggcg	180
74	gcctcagtgaa gcgacgcgc ggcagagag ggagtggcca actccatcac taggggttc	240
75	tttgttagttaa tgatataaccg gccatgtctac ttatctaccg atgatgtca gcttgcac	300
77	<210> SEQ ID NO: 6	
78	<211> LENGTH: 272	
79	<212> TYPE: DNA	
80	<213> ORGANISM: AAV circular intermediate, clone p1202	
82	<400> SEQUENCE: 6	
83	gcatcaagc ttagataag tagcatggcg ggttaatcat taactacaag gaacccttag	60
84	ttagggatt ggccactccc tctctgcgcg ctgcgtcgct cactgaggcc gggcgaccaa	120
85	aggtcggccg acggccggc ttgggtcgcc cggcgtcgat ggcgacgcgaa ggcgcagag	180
86	agggagtggc caactccatc actaggggtt ctttgttagtt aatgattaac ccgcctatgc	240
87	acttatctac cgatgaattc gagcttgcac gc	272
89	<210> SEQ ID NO: 7	
90	<211> LENGTH: 165	
91	<212> TYPE: DNA	
92	<213> ORGANISM: Unknown	
94	<220> FEATURE:	
95	<223> OTHER INFORMATION: SEQ ID NO:1 of U.S. Patent No. 5,478,745	
97	<400> SEQUENCE: 7	
98	aggaacccct agtgatggg ttggccactc cctctctgcg cgctcgctcg ctcaactgagg	60
99	ccggcgacc aaaggctgcc cgaegccgg gctttgccc ggcggcctca gtgagcgac	120
100	gagcgcgcag agaggagtg gccaactcca tcactagggg ttcc	165
102	<210> SEQ ID NO: 8	
103	<211> LENGTH: 282	
104	<212> TYPE: DNA	
105	<213> ORGANISM: rAAV circular intermediate, clone p79	
107	<400> SEQUENCE: 8	
108	ggcgccat ttatcgtaag ttatgtggcg actgcaggca tgcaagctcg aattcatcg	60
109	tagatagtt gcatggcggtt ttaatcattt cttacaaaaa gcccctatgt atggatggg	120
110	ccactccctc ttttcgcgcg ggcgcgcagag agggatggc caactccctc actaggggtt	180
111	cctggcgtt aatgatgtac ccgcctatgc acttatctac agttgcacat catgtgagca	240
112	aaaggccagaaaaggccag gaaccgtaaa aaggccgcgt tg	282
114	<210> SEQ ID NO: 9	
115	<211> LENGTH: 345	
116	<212> TYPE: DNA	
117	<213> ORGANISM: rAAV circular intermediate, clone p80	
119	<400> SEQUENCE: 9	
120	ggccattac cgtaatgtt gtaacgactg caggcatgca agtcgaatt catggtaga	60
121	taatgtcat ggcgggttaa tcattacta caaggaaccc ctatgtatgg agttggccac	120

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122	tccctctcg	cgcgtcgct	cgtcgctca	ggccggcga	ccaaaggcg	cccacgccc	180
123	gccccgctc	agcgagcgag	cgagccgcga	gagaggaga	ggccaactcc	atca	240
124	gttcctgt	gttaatgatt	aaccgcct	gctacttac	tacagcttg	atgc	300
125	gcaaaaaggc	agcaaaaaggc	caggaaccgt	aaaaaggcg	cgttg		345
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128	<211>	LENGTH:	276				
129	<212>	TYPE:	DNA				
130	<213>	ORGANISM:	rAAV circular intermediate, clone p81				
132	<400>	SEQUENCE:	10				
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134	taatgtat	ggcggtttaa	tcaatgccta	caaagagcc	ctagtatgg	agccggcct	120
135	caccgagcga	gcgagcgcgc	agagaggag	tggccaaactc	catca	ctagg ggttcc	180
136	agttatgtat	taaccgcct	tgcata	ttt	atcagcttg	catgc	240
137	cagcaaaaagg	ccaggaaccg	taaaaaggcc	gcgtt			276
139	<210>	SEQ ID NO:	11				
140	<211>	LENGTH:	316				
141	<212>	TYPE:	DNA				
142	<213>	ORGANISM:	rAAV circular intermediate, clone p86				
144	<400>	SEQUENCE:	11				
145	ggccattttac	cgttaagtat	gtaaacgactg	caggcatgca	agctcgaa	att catcggt	60
146	taatgtat	ggcggtttaa	tcaatacta	caaggaaccc	ctagtatgg	agttggcac	120
147	tccctctcg	cgcgtcgct	cgtcgctga	ggccggcccg	gcctcagcga	gcgagcgcgc	180
148	gcgagagag	ggactggcca	actccatcac	taggggttcc	ttgtat	ttaa tgattaaccc	240
149	gccatgtat	tatctacag	cttgc	catgca	tgttagcaaa	aggccagcaa	300
150	accgtaaaaa	ggccgc					316
152	<210>	SEQ ID NO:	12				
153	<211>	LENGTH:	208				
154	<212>	TYPE:	DNA				
155	<213>	ORGANISM:	rAAV circular intermediate, clone p87				
157	<400>	SEQUENCE:	12				
158	ggccattttac	cgttaagtat	gtaaacgactg	caggcatgca	agctcgaa	att catcggt	60
159	taatgtat	ggcggtttaa	tcaatgccta	caaagagcc	ctagtatgg	aattggat	120
160	attcacccctc	catgtatctt	atctacagct	tgc	atgcatg	ttagcaaa	180
161	ggccaggaac	cgtaaaaagg	ccgcgtt				208
163	<210>	SEQ ID NO:	13				
164	<211>	LENGTH:	310				
165	<212>	TYPE:	DNA				
166	<213>	ORGANISM:	rAAV circular intermediate, clone p88				
168	<400>	SEQUENCE:	13				
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170	aatgtat	ggcggtttaa	cattgcctac	aaagagcc	tagtatgg	gttggccact	120
171	tccctctcg	cgcgtcgctc	gctggggcccg	gcctcagcga	gcgagcgcgc	gcccagagag	180
172	ggatggcca	actccatcac	taggggttcc	ttgtat	ttaa tgattaaccc	gccatgtac	240
173	ttatctacag	cttgc	atgca	tgttagcaaa	aggccagcaa	aaggccagga	300
174	ggccgcgtt						310

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L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:39 M:283 W: Missing Blank Line separator, <220> field identifier